## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/56/902
Source:	1.FWP
Date Processed by STIC:	1/3/06

## ENTERED



**IFWP** 

RAW SEQUENCE LISTING DATE: 01/03/2006
PATENT APPLICATION: US/10/561,902 TIME: 11:33:18

Input Set : N:\RJAVED\10561902.txt
Output Set: N:\CRF4\01032006\J561902.raw

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3 <110> APPLICANT: Universitatsklinikum Freiburg
 5 <120> TITLE OF INVENTION: Modulation of Osteoblast Activity by Fhl2
 7 <130> FILE REFERENCE: Fhl2
9 <140> CURRENT APPLICATION NUMBER: US/10/561,902
9 <141> CURRENT FILING DATE: 2005-12-21
 9 <160> NUMBER OF SEQ ID NOS: 3
11 <170> SOFTWARE: PatentIn version 3.1
13 <210> SEQ ID NO: 1
14 <211> LENGTH: 840
15 <212> TYPE: DNA
16 <213> ORGANISM: homo sapiens
18 <400> SEQUENCE: 1
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21 ctgcgggagg agagccccta ctgcgtggtg tgctttgaga ccctgttcgc caacacctgc
                                                                         120
23 gaggagtgtg ggaagcccat cggctgtgac tgcaaggact tgtcttacaa ggaccggcac
                                                                         180
25 tggcatgaag cctgtttcca ctgctcgcag tgcagaaact cactggtgga caagcccttt
                                                                         240
27 gctgccaagg aggaccagct gctctgtaca gactgctatt ccaacgagta ctcatccaag
                                                                         300
29 tgccaggaat gcaagaagac catcatgcca ggtacccgca agatggagta caagggcagc
                                                                         360
31 agetggeatg agacetgett catetgeeae egetgeeage agecaattgg aaceaagagt
                                                                         420
33 ttcatcccca aagacaatca gaatttctgt gtgccctgct atgagaaaca acatgccatg
                                                                         480
                                                                         540
35 caqtqcqttc aqtqcaaaaa qcccatcacc acqqqaqqqq tcacttaccg ggagcagccc
37 tggcacaagg agtgcttcgt gtgcaccgcc tgcaggaagc agctgtctgg gcagcgcttc
                                                                         600
39 acagetegeg atgaetttge ctaetgeetg aactgettet gtgaettgta tgecaagaag
                                                                         660
41 tgtgctgggt gcaccaaccc catcagcgga cttggtggca caaaatacat ctcctttgag
                                                                         720
                                                                         780
43 qaacqqcaqt qqcataacqa ctqctttaac tqtaaqaaqt gctccctctc actggtgggg
45 cgtggcttcc tcacagagag ggacgacatc ctgtgccccg actgtgggaa agacatctga
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48 <210> SEQ ID NO: 2
49 <211> LENGTH: 279
50 <212> TYPE: PRT
51 <213> ORGANISM: homo sapiens
53 <400> SEQUENCE: 2
55 Met Thr Glu Arg Phe Asp Cys His His Cys Asn Glu Ser Leu Phe Gly
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56 1
59 Lys Lys Tyr Ile Leu Arg Glu Glu Ser Pro Tyr Cys Val Val Cys Phe
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60
               20
63 Glu Thr Leu Phe Ala Asn Thr Cys Glu Glu Cys Gly Lys Pro Ile Gly
64
           35
                                40
67 Cys Asp Cys Lys Asp Leu Ser Tyr Lys Asp Arg His Trp His Glu Ala
71 Cys Phe His Cys Ser Gln Cys Arg Asn Ser Leu Val Asp Lys Pro Phe
                       70
                                            75
75 Ala Ala Lys Glu Asp Gln Leu Leu Cys Thr Asp Cys Tyr Ser Asn Glu
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79 Tyr Ser Ser Lys Cys Gln Glu Cys Lys Lys Thr Ile Met Pro Gly Thr
80
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83 Arg Lys Met Glu Tyr Lys Gly Ser Ser Trp His Glu Thr Cys Phe Ile
                               120
84
           115
                                                    125
87 Cys His Arg Cys Gln Gln Pro Ile Gly Thr Lys Ser Phe Ile Pro Lys
                           135
91 Asp Asn Gln Asn Phe Cys Val Pro Cys Tyr Glu Lys Gln His Ala Met
                       150
                                            155
95 Gln Cys Val Gln Cys Lys Lys Pro Ile Thr Thr Gly Gly Val Thr Tyr
96
                   165
                                       170
99 Arq Glu Gln Pro Trp His Lys Glu Cys Phe Val Cys Thr Ala Cys Arg
                                    185
103 Lys Gln Leu Ser Gly Gln Arg Phe Thr Ala Arg Asp Asp Phe Ala Tyr
104
            195
                                200
107 Cys Leu Asn Cys Phe Cys Asp Leu Tyr Ala Lys Lys Cys Ala Gly Cys
111 Thr Asn Pro Ile Ser Gly Leu Gly Gly Thr Lys Tyr Ile Ser Phe Glu
112 225
                        230
                                             235
115 Glu Arg Gln Trp His Asn Asp Cys Phe Asn Cys Lys Lys Cys Ser Leu
                    245
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119 Ser Leu Val Gly Arg Gly Phe Leu Thr Glu Arg Asp Ile Leu Cys
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123 Pro Asp Cys Gly Lys Asp Ile
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127 <210> SEQ ID NO: 3
128 <211> LENGTH: 1892
129 <212> TYPE: DNA
130 <213> ORGANISM: homo sapiens
132 <400> SEQUENCE: 3
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135 ggcgcccgcc ccccgcgccc cacgcccggt gccagcgagc cgaggcgtgc atctccttat
                                                                          120
137 atggtcaaat gacacggcgg ggtttctcga gggcgggagc tgcgcagcgc tccactcggc
                                                                          180
                                                                          240
139 eggeagegga geegeageea ceageegeee gegeeeteea geecegteeg ggagteeeeg
141 gcccqctqcq qtqccqtqaq tacctccaac cccctqcgcc ccgqagggag gccgaggggc
                                                                          300
                                                                          360
143 ttagccacca gggctcggaa gtgggggccg aatccggtgc gagacccaag gagaggggag
                                                                          420
145 cagageegga gttggggaga etgtggetga aaactgtgte tteetggaga etaggetgge
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147 attttgactt tgggacggag tetegetttg tegeccagge tggagtgeag tggcacgate
149 tcagctcact gcaagctcta cctcttggtt cacgccattc tcctgcccca gcctcccaag
                                                                          540
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151 tagctgggac tacaggttgc tgaaaagcca ggagtcaaaa tgactgagcg ctttgactgc
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153 caccattgca acgaatctct ctttggcaag aagtacatcc tgcgggagga gagcccctac
                                                                          720
155 tgcgtggtgt gctttgagac cctgttcgcc aacacctgcg aggagtgtgg gaagcccatc
157 ggctgtgact gcaaggactt gtcttacaag gaccggcact ggcatgaagc ctgtttccac
                                                                          780
159 tgctcgcagt gcagaaactc actggtggac aagccctttg ctgccaagga ggaccagctg
                                                                          840
                                                                          900
161 ctctgtacag actgctattc caacgagtac tcatccaagt gccaggaatg caagaagacc
163 atcatgccag gtacccgcaa gatggagtac aagggcagca gctggcatga gacctgcttc
                                                                          960
165 atctgccacc gctgccagca gccaattgga accaagagtt tcatccccaa agacaatcag
                                                                         1020
                                                                         1080
167 aatttetgtg tgeectgeta tgagaaacaa catgeeatge agtgegttea gtgeaaaaag
169 cccatcacca cgggagggt cacttaccgg gagcagccct ggcacaagga gtgcttcgtg
                                                                         1140
171 tgcaccgcct gcaggaagca gctgtctggg cagcgcttca cagctcgcga tgactttgcc
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173	tactgcctga	actgcttctg	tgacttgtat	gccaagaagt	gtgctgggtg	caccaacccc	1260
175	atcagcggac	ttggtggcac	aaaatacatc	tcctttgagg	aacggcagtg	gcataacgac	1320
177	tgctttaact	gtaagaagtg	ctccctctca	ctggtggggc	gtggcttcct	cacagagagg	1380
179	gacgacatcc	tgtgccccga	ctgtgggaaa	gacatctgaa	ttcaacacag	agaagttgct	1440
181	gcttgtgatc	tcacacacag	atttttatgt	tttctttctc	acccaggcaa	tcttgccttc	1500
183	tggtttcttc	cagccacatt	gagactttct	tctagtgctt	ttcagtgata	ctcacgtttg	1560
185	cttaaaccct	ttagtgcttt	gtgatagttc	agtcccaggg	aaagagaaaa	ctcgccctag	1620
187	gccctaggtg	ggaagatggt	ttgaaatttt	tgtaatcgag	taaggcacac	ccaaatgtaa	1680
189	aaatcctttt	gaatgatgcc	tttataaatc	tttctctcac	tgtctattta	agtgcaatta	1740
191	acatatgtca	cgaacttgaa	agttttctaa	actcaataag	gtaatgacca	gttgttattt	1800
193	acagctctgt	aacctcccgt	tgcgtcaagt	ctaaaccaag	attatgtgac	ttgcaataaa	1860
195	gttattcaga	acagaaaaaa	aaaaaaaaa	aa			1892

VERIFICATION SUMMARY

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L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date